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B

PCR Fragg./ CDNA clones	nt. positions
PCR1	388-731
PCR2	1575-1774
PCR3	388-1774
5'RACE	1-540
G6 library	1-641
1.1	53-1007
1.2	1018-2188
1.3	1553-2563
2.1	451-3837
2.2	1590-3983
2.3	2315-3983

A

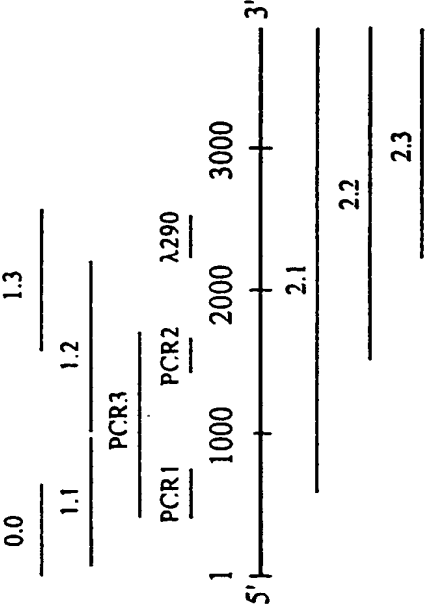


Fig. 1

SUBSTITUTE SHEET (RULE 26)

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Fig 2b

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L K E F P E E L K N H G A Y L G Y T V T S V V S S R Q G R V Y V A G A P R F N H 450

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T G K V I L F T M H N N R S L T I H Q A H R G Q Q I G S Y F G S E I T S V D I D 490

GGGAGCGGCTGACTGATGCTCTGCTGGTGGCGCACCCATGTACTTCAACGAGGGCGGTGAGCGAGGCAAGGTGTACCTCTATGAGCTGAGACAGAACCGGTTTGTATTATAACGGAACG 1680
G D G V T D V L L V G A P H Y F N E G R E R G K V Y Y E L R Q N R F V Y N G T 530

CTAAGGATTACACAGTTACCAAGATGCCGATTTGGGTCTCTCCATTGCTCAGTTCCAGACTCAACAGGATTCCTACATGCTACATGCTGGTGGAGCGCCCTGGAGGACCAACCAC 1800
L K D S H S Y Q N A R F G S S I A S V R D L N O D S Y N D V V V G A P L E D N H 570

GCAGGAGCCATCTACATCTTCCAGGCTTCCGAGGAGCATCTCTGAAGACACCTAAGCAGAGATCACAGCCTCAGAGCTGGCTACCGGCTCCAGTATTTTGGCTGCAGCATCCACGGG 1920
A G A I Y I F H G F R G S I L K T P K Q R I T A S E L A T G L Q Y F G C S I H G 610

CAATTGGACCTCAATGAGGATGGCTCATCGACCTGGCAGTGGGAGCCCTTGGCAGCTGTGATTTCTGTGGTCCCGCCAGTGGTTTCAGATCAATGCCAGCCTCCACTTTGAGCCATCC 2040
Q L D L N F D G L I D L A V G A L G N A V I L W S R P V V Q I N A S L H F E P S 650

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K I N I F H R D C K R S G R D A T C L A A F L C F T P I F L A P H F Q T T T V G 690

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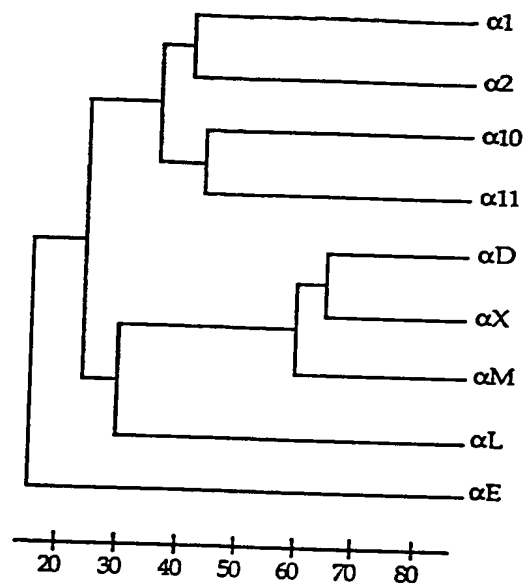
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Fig 2c

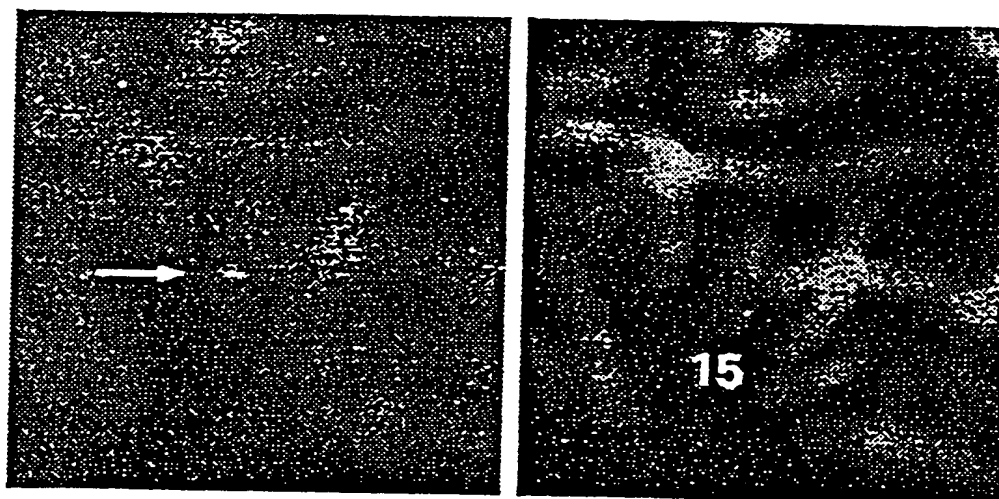
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 I A T R S G N R L L K L R D F L T D E A N[•] T S C N I W G N[•] S T E Y R P T P V E E 1050
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 GCCCCAGCTACTAAGGTGCTAGG 3983

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**Fig. 3**

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A



B

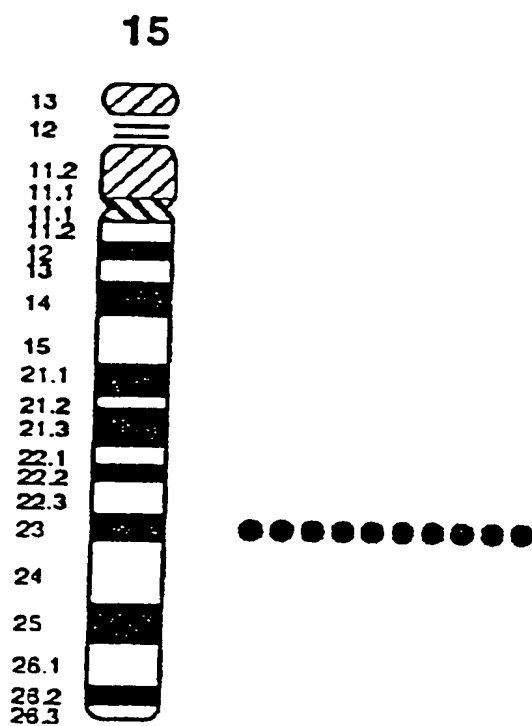


FIG. 4

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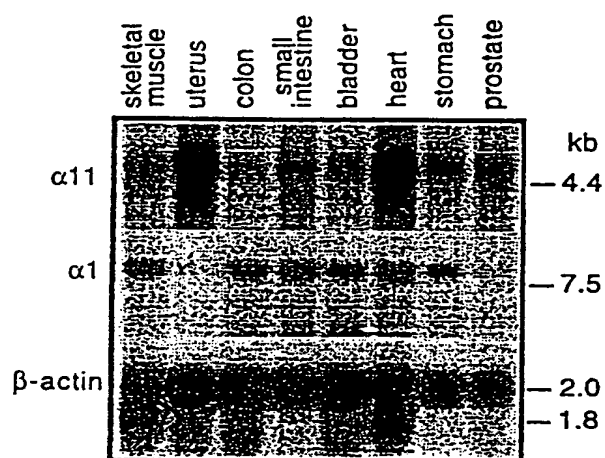


Fig. 5

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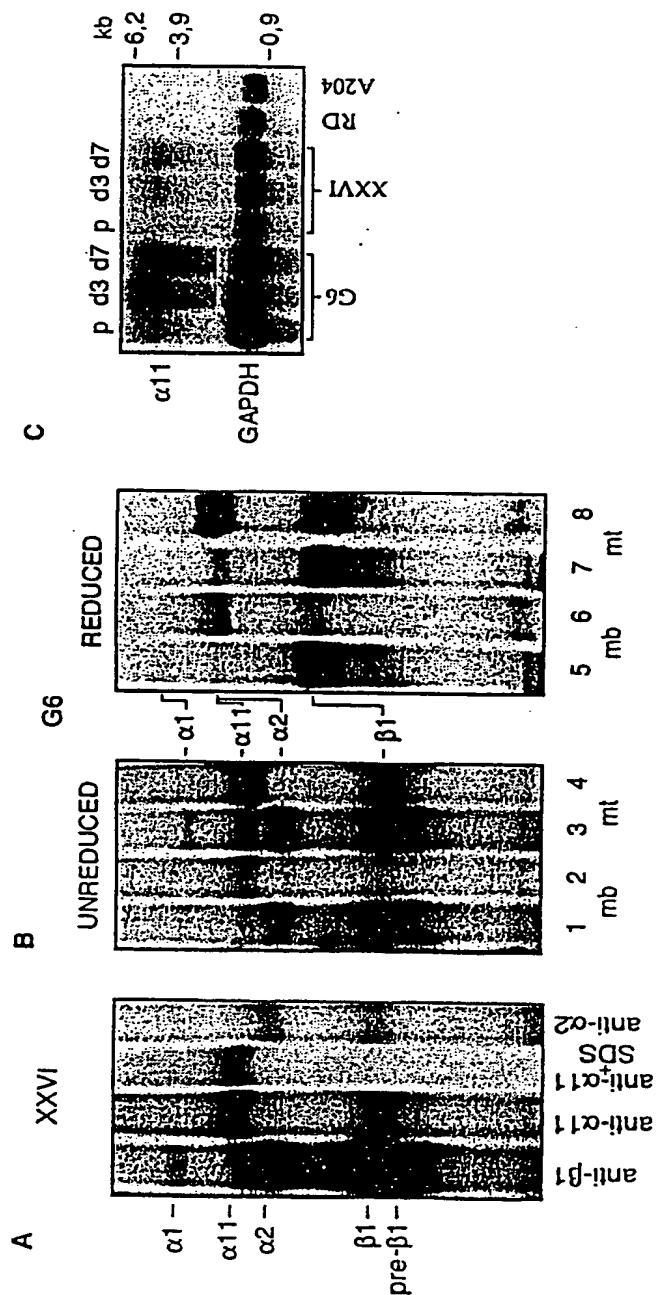


Fig. 6

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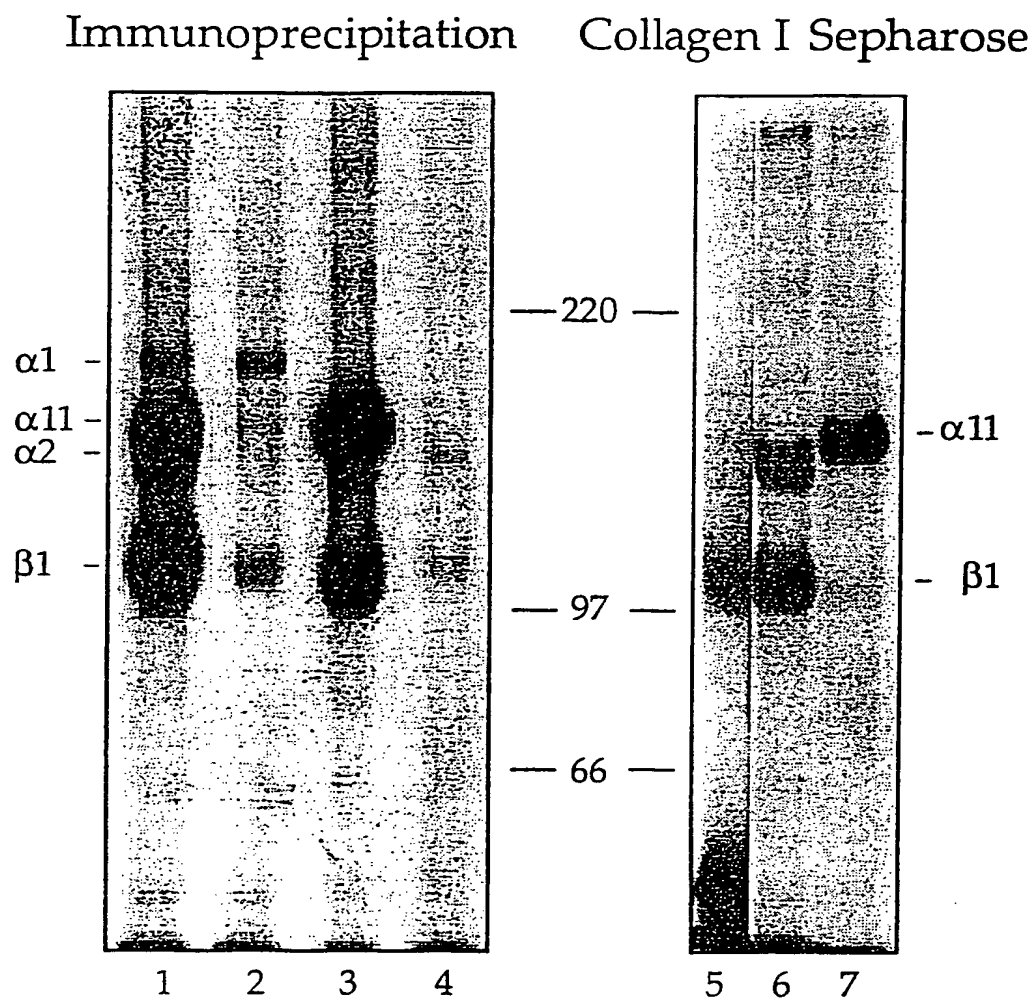


Fig. 7 panel 1

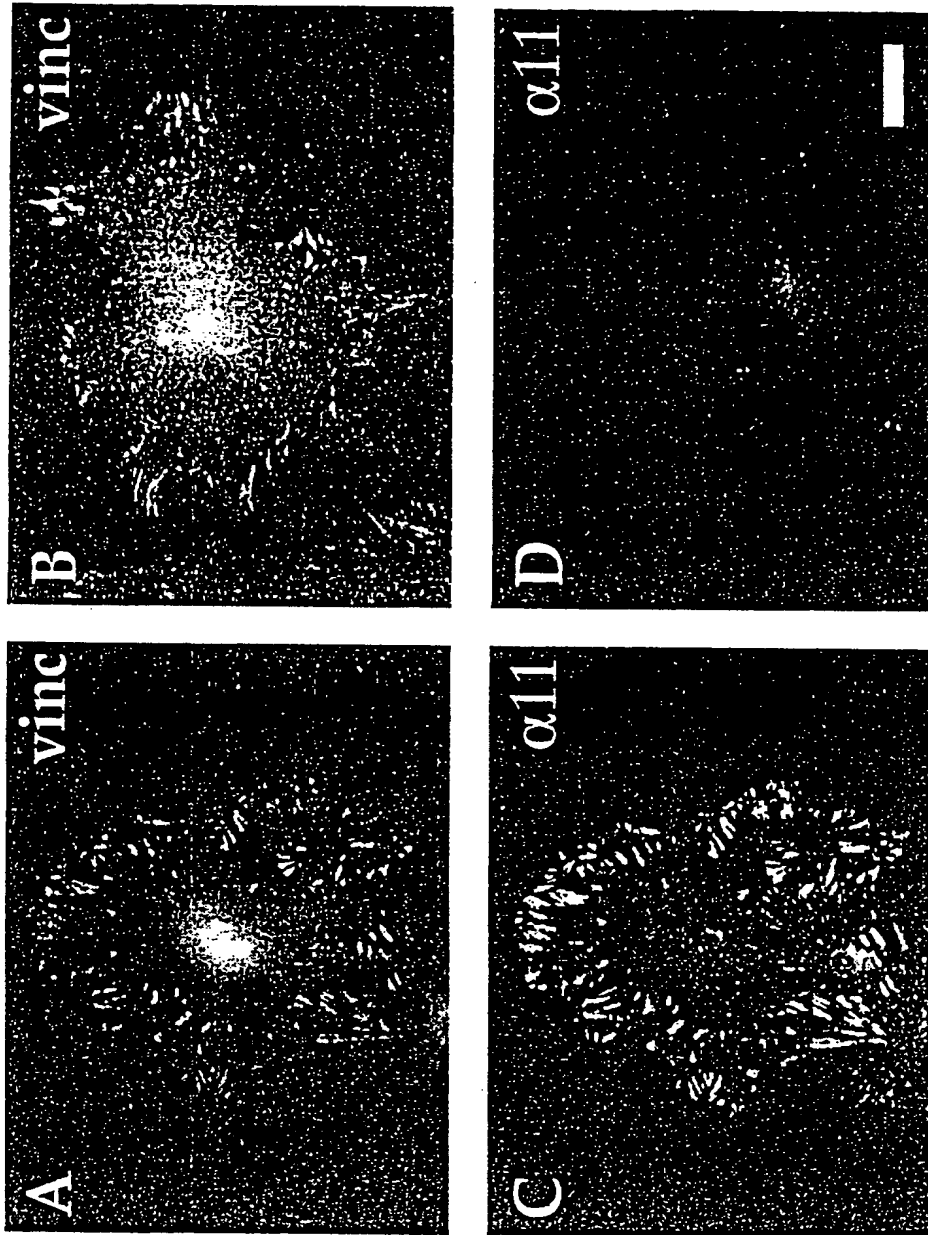


Fig. 7 panel 2

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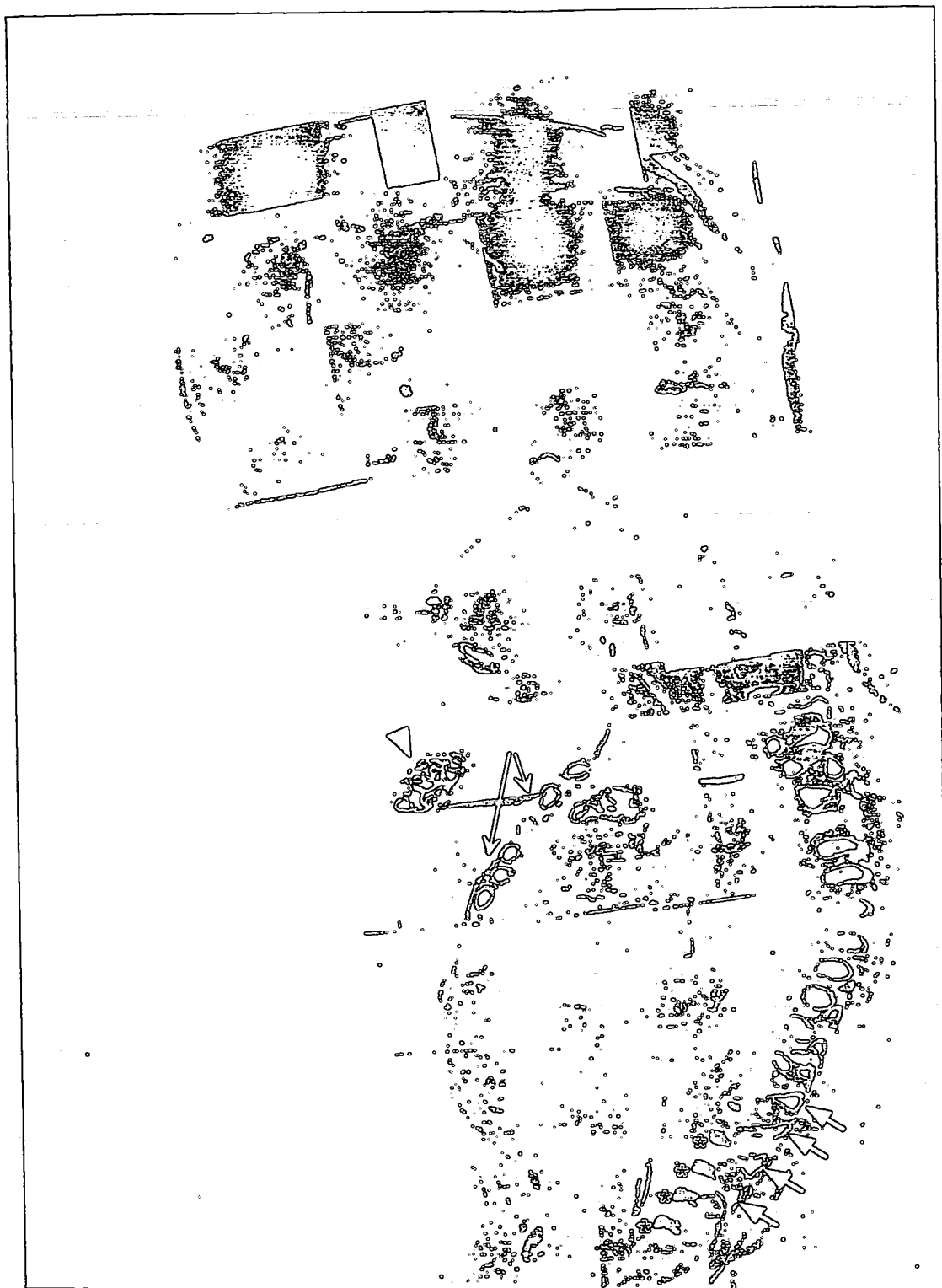


FIG. 8